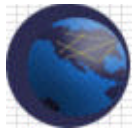


**Integrative Cancer Research Workspace**

**Data Analysis and Statistical Tools**

**Special Interest Group Teleconference**

Date, Time & Location:	November 5, 2004 2:00-3:00 PM EST		
Attendees:			
	First Name	Last Name	Institution
	Ted	Liefeld	Broad/MIT
	Wen-Pin	Chen	Chao Family Comprehensive Cancer Center
	Bob	Clarke	Georgetown University
	Jason	Xuan	Georgetown University
	Subha	Madhavan	National Cancer Institute
	David	Kane	NCI CCR
	Julie	Zhu	Northwestern University
	Ajay	Jain	UCSF
	Everett	Zhou	UNC
	Chris	Kingsley	University of California
	Xiaoning	Wang	University of Chicago
	Leo	Cheung	University of Hawaii
	Lynne	Wilkins	University of Hawaii
Juli	Klemm	3rd Millennium	
Presentation	<b><u>Magellan Presentation</u></b>		
	Chris Kingsley (UCSF) gave a presentation of Magellan. The slides can be downloaded from <a href="http://cabig.nci.nih.gov/workspaces/ICR/Meetings/SIGs/Data_analysis_stats/20041105_Magellan_presentation/view">http://cabig.nci.nih.gov/workspaces/ICR/Meetings/SIGs/Data_analysis_stats/20041105_Magellan_presentation/view</a>		
Discussion	<b><u>Support from Cross-cutting Workspaces:</u></b>		
	List of architecture mentors for the projects in this SIG was emailed on 11/2/04 and can be downloaded from: <a href="http://cabig.nci.nih.gov/workspaces/ICR/Points_of_Contact/ICR_SIG_contactlist_20041110">http://cabig.nci.nih.gov/workspaces/ICR/Points_of_Contact/ICR_SIG_contactlist_20041110</a>		
	<b><u>Integration of DA/ST projects with caBIG</u></b>		
	Initiated a discussion among project representatives on the topic of data representation, object models, vocabularies, etc. Heard from reps of DWD and GenePattern.  Steve Marron (DWD) brought up the question of how best to produce MAGE-ML from data in the Stanford format. It was suggested that caArray can produce MAGE-ML and that data can be loaded in using the EJB API that is read/write. It was also suggested that the caArray API could be used read data from caArray using the MAGE-OM API.  Discussions on this topic to be continued in future meetings as developers move toward integration of their applications with caBIG (possibly focusing on the acquisition of data and annotations from caArray).		



**caBIG** *cancer Biomedical  
Informatics Grid*



Action Items:	Name Responsible	Action Item	Date Due	Notes
	Chris Kingsley	Distribute meeting minutes	11/15/04	
	Juli Klemm	Post Presentation Slides on caBIG website	11/10/04	